

GenCore version 4.5  
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Om protein - protein search, using sw model

Run on: March 1, 2001, 15:47:15 ; Search time 210.42 Seconds  
(without alignments)  
6.988 Million cell updates/sec

Title: US-09-331-631A-5\_COPY\_33\_75  
 Perfect score: 340

Sequence: 1 NQEDPQTECQCCQRCRQDE.....RQQQYCCRRCKEICEEEFY 43

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 268485 seqs, 34193795 residues

Total number of hits satisfying chosen parameters: 268485

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

A\_Geneseq\_36: \*

1:	/SIDS1/gcgdata/geneseq/geneseqp/AA1980.DAT *
2:	/SIDS1/gcgdata/geneseq/geneseqp/AA1981.DAT *
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6:	/SIDS1/gcgdata/geneseq/geneseqp/AA1985.DAT *
7:	/SIDS1/gcgdata/geneseq/geneseqp/AA1986.DAT *
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13:	/SIDS1/gcgdata/geneseq/geneseqp/AA1992.DAT *
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16:	/SIDS1/gcgdata/geneseq/geneseqp/AA1995.DAT *
17:	/SIDS1/gcgdata/geneseq/geneseqp/AA1996.DAT *
18:	/SIDS1/gcgdata/geneseq/geneseqp/AA1997.DAT *
19:	/SIDS1/gcgdata/geneseq/geneseqp/AA1998.DAT *
20:	/SIDS1/gcgdata/geneseq/geneseqp/AA1999.DAT *
21:	/SIDS1/gcgdata/geneseq/geneseqp/AA2000.DAT *

**Pred. No.** is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

## SUMMARIES

Result	Score	Query	Match	Length	DB	ID	Description
No.							
1	248	100.0		625	19	W62830	Macadamia integrifolia
2	241	97.2		666	19	W62828	Macadamia integrifolia
3	235	94.8		666	19	W62839	Macadamia integrifolia
4	110	44.4		525	19	W62831	Theobroma cacao an
5	110	44.4		566	13	R20181	Sequence encoded b
6	109	44.0		590	19	W62832	Gossypium hirsutum
7	96	38.7		28	19	W62841	Stenocarpus sinuata
8	68.5	27.6		593	19	W62835	Zea mays antilmicro
9	66.5	26.8		35	13	R21079	Zea mays antilmicro
10	66.5	26.8		637	19	W62837	Hordeum vulgare an
11	65.5	26.4		33	19	W62836	Zea mays antilmicro
12	65	26.2		910	20	Y22191	Mouse brain CNG-1

13	65	26.2	919	18	P93109	Human androgen receptor
14	65	26.2	919	18	M14783	Androgen receptor
15	65	26.2	919	21	T78914	Human androgen receptor
16	63.5	25.6	51	18	M33694	Mouse prolactin 1
17	63.5	25.6	176	18	M33695	Mouse prolactin 1
18	63.5	25.6	301	19	M37085	Anti-human SC.sing
19	63.5	25.6	409	20	W60342	G. max truncated S
20	63.5	25.6	489	20	M90341	G. max SBP2 protease
21	63	25.4	626	18	M22150	Peanut allergen Ar
22	63	25.4	626	20	Y15244	Peanut allergen Ar
23	63	25.4	626	20	Y25657	Peanut allergen Ar
24	62	25.0	1162	21	Y58500	Human 5. EST relat
25	61.5	24.8	98	21	Y65429	Human cyclin D3 ps
26	61.5	24.8	215	14	R44806	Murine PCIP protea
27	61	24.6	1447	20	M81029	Human unliganded r
28	60	24.2	154	20	Y33504	Human androgen rec
29	60	24.2	918	12	R12323	Peanut allergen Ar
30	60	24.2	918	20	Y33491	Arachis hypogaea A
31	59.5	24.0	614	18	W22149	Amino acid sequenc
32	59.5	24.0	614	19	W62324	Sequence of human
33	59	23.8	2023	21	Y54320	A secreted protein
34	58.5	23.6	461	15	R51002	Neisseria meningit
35	58.5	23.6	514	19	M80400	Neisseria meningit
36	58	23.4	151	21	Y74634	Human EDF-binding
37	58	23.4	199	21	Y74635	Human follistatin
38	57.5	23.2	304	13	R20063	Glycine max antilm
39	57.5	23.2	317	10	P93396	Human cerebral pro
40	57.5	23.2	605	19	W62338	Diofilaria immitis
41	57.5	23.0	809	20	Y29672	A human trichonyal
42	57	23.0	303	15	R60054	Amino acid sequenc
43	57	23.0	1898	20	Y30795	Human cytoskeleton
44	57	23.0	2074	21	Y54319	
45	56.5	22.8	281	21	Y91958	

## ALIGNMENTS

RESULT	1
W62830	
ID	W62830 standard; Protein; 625 AA.
XX	
AC	W62830;
XX	
DT	27-OCT-1998 (first entry)
XX	
DE	Macadamia integrifolia antimicrobial protein.
XX	
KW	antimicrobial protein; infestation; control.
XX	
OS	Macadamia integrifolia.
XX	
Key	Location/Qualifiers
FT	1..28
FT	/note="signal peptide"
FT	29..666
FT	/note="mature protein"
XX	
PN	W09827805-A1.
XX	
PD	02-JUL-1998.
XX	
PF	22-DEC-1997; 97MO-AU00874.
XX	
PR	20-DEC-1996; 96AU-0004275.
XX	
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX	
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP
DR	WPI: 1998-377279/32.
DR	N-PSDB; V42316.
XX	

```
PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
PT useful for controlling microbial infestations of plants or mammals
XX
PS Claim 1; Page 43-45; 96pp; English.
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 625 AA;

Query Match 100.0%; Score 248; DB 19; Length 625;
Best Local Similarity 100.0%; Pred. No. 1.3e-19;
Matches 43; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 NOEDPQTECQOCQRRRCROESDPROOQYCORCKEICEEEY 43
Db 33 ngedpqtceqgcqrrrcrqsqsdprqgycqrrckelceeeey 75

RESULT 2
W62828 standard; Protein; 666 AA.
AC W62828;
DT 27-OCT-1998 (first entry)
DE Macadamia integrifolia antimicrobial protein.
KW antimicrobial protein; infestation; control.
OS Macadamia integrifolia.
XX
XX Key Location/Qualifiers
FH Peptide 1..28
FT /note= "signal peptide"
FT 29..666
FT Protein /note= "mature protein"
PN W09827805-A1.
PD 02-JUL-1998.
XX
XX 22-DEC-1997; 97WO-AU00874.
XX
XX 20-DEC-1996; 96AU-0004275.
XX
PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX
PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
DR WPI: 1998-377279/32.
DR N-PSDB; VA2310.
XX
XX Novel anti-microbial protein from e.g. Macadamia integrifolia -
PT useful for controlling microbial infestations of plants or mammals
XX
PS Claim 1; Page 34-36; 96pp; English.
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 666 AA;

Query Match 97.2%; Score 241; DB 19; Length 666;
Best Local Similarity 97.7%; Pred. No. 7.8e-19;
Matches 42; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 NOEDPQTECQOCQRRRCROESDPROOQYCORCKEICEEEY 43
```

```
Db 74 ngedpqtceqgcqrrrcrqsqsdprqgycqrrckelceeeey 116

RESULT 3
W62829 standard; Protein; 666 AA.
AC W62829;
DT 27-OCT-1998 (first entry)
DE Macadamia integrifolia antimicrobial protein.
KW antimicrobial protein; infestation; control.
OS Macadamia integrifolia.
XX
XX Key Location/Qualifiers
FH Peptide 1..28
FT /note= "signal peptide"
FT 29..666
FT Protein /note= "mature protein"
PN W09827805-A1.
PD 02-JUL-1998.
XX
XX 22-DEC-1997; 97WO-AU00874.
XX
XX 20-DEC-1996; 96AU-0004275.
XX
PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX
PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
DR WPI: 1998-377279/32.
DR N-PSDB; VA2311.
XX
XX Novel anti-microbial protein from e.g. Macadamia integrifolia -
PT useful for controlling microbial infestations of plants or mammals
XX
PS Claim 1; Page 39-41; 96pp; English.
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 666 AA;

Query Match 94.8%; Score 235; DB 19; Length 666;
Best Local Similarity 93.0%; Pred. No. 3.5e-18;
Matches 40; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 NOEDPQTECQOCQRRRCROESDPROOQYCORCKEICEEEY 43
Db 74 ngedpqtceqgcqrrrcrqsqsdprqgycqrrckelceeeey 116

RESULT 4
W62831 standard; Protein; 525 AA.
AC W62831;
DT 27-OCT-1998 (first entry)
DE Theobroma cacao antimicrobial protein.
KW antimicrobial protein; infestation; control.
OS Theobroma cacao.
```

XX	MO9827805-A1.
PN	
XX	
XX	02-JUL-1998.
PD	
XX	
PF	22-DEC-1997; 97MO-AU00874.
XX	
PR	20-DEC-1996; 96AU-0004275.
XX	
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX	
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
DR	WPI; 1998-377279/32.
XX	
PT	Novel anti-microbial protein from e.g. <i>Macadamia integrifolia</i> -
XX	useful for controlling microbial infestations of plants or mammal
PS	Claim 1; Page 47-49; 96pp; English.
XX	
CC	The sequence is that of an antimicrobial protein which can
XX	be used to control microbial infestations in plants and mammalian
XX	animals.
XQ	Sequence 525 AA;

```

Oxy Match 19: 44.4% Score 110; DB 19: Length 525;
Best Local Similarity 47.5%; Pred. No. 9,8e-05;
Matches 19: Conservative 10; Mismatches 11; Indels 0; Gaps 0
OY 2 QEDPTECCGCCRCRQESDPRQOYTCRRCKICEEE 41
    . . . . . 1111 11:1: 1:11 111:1: 1:1
Db 78 eeelqrayqcgqrcgqgqgqgqrcgkewgykce 117

```

CC	XX	RESULT	5
CC	XX	ID	R20181
CC	XX	R20181	standard; Protein; 566 AA.
CC	XX	R20181;	
CC	XX	DT	16-APR-1992 (first entry)
CC	XX	DE	Sequence encoded by 67 kD T. cacao protein cDNA.
CC	XX	KM	Cocoa; flavour; vicilin; seed storage protein.
CC	XX	OS	Theobroma cacao.
CC	XX	PN	WO9119801-A.
CC	XX	PD	26-DEC-1991.
CC	XX	PF	07-JUN-1991; 91WO-GB000914.
CC	XX	PR	11-JUN-1990; 90GB-0013016.
CC	XX	PA	(MRS C ) MARS UK LTD.
CC	XX	PI	Spencer ME, Hodge R, Deakin EA, Ashton S;
CC	XX	DR	WPI; 1992-024418/03.
CC	XX	DR	N-PDB; Q20377.
CC	XX	PT	Recombinant cocoa proteins - are responsible for flavour in cocoa
CC	XX	PT	beans and produced in large quantities using yeast and bacterial
CC	XX	PS	expression vectors
CC	XX	PS	Claim 4; Fig 2; 59pp: English.
CC	XX	CC	The inventors claim a 67 kD and 31 kD T. cacao protein, and
CC	XX	CC	fragments, and encoding DNAs. The 47 kD and 31 kD proteins are

CC derived from the 67 kD precursor. T. cacao protein cDNA was  
CC detected in a cDNA library prepared from immature cocoa beans RNA  
CC using a probe based on the Aa sequence of a CNBr peptide common to  
CC the 47 kD and 31 kD polypeptides. Homology searches revealed close  
CC homologies between the 67 kD polypeptide and the vicilins, which are  
CC seed storage proteins.

XX Sequence 566 AA;  
50

Query Match	44.4%	Score 110	DB 13	Length 566
Best Local Similarity	47.5%	Pred. No.	0.0001	
Matches 19	Conservative 10	Mismatches 11	Indels 0	Gaps 0

```

QY      2 QDEPQTFEGQCCQRCRCRQDESPDQQQYQRCRKEICIEEEE 41
      :: : : ||| ||::: : ||| ||| : : ::|
Db      78 eeelgrqyqgcgrcgcgcqgqgcqgcqgcqrkckweykeke 117

```

RESULT	6
W62832	
ID	W62832 standard; Protein; 590 AA.
XX	
AC	W62832:
XX	
DT	27-OCT-1998 (first entry)
XX	
DE	Gossypium hirsutum antimicrobial protein.
XX	
DE	antimicrobial protein; infestation; control
KW	

XX	Gossypium hirsutum.
XX PN	M09827805-A1.
XX PD	02-JUL-1998.
XX PF	22-DEC-1997; 97WO-AU00874.
XX PR	20-DEC-1996; 96AU-0004275.
XX PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX DR	WPI; 1998-377279/32.
PT	Novel antimicrobial protein from e.g. Macadamia integrifolia -
PT	useful for controlling microbial infestations of plants or mammals
PS PS	Claim 1; Page 49-51; 96pp; English.
CC CC	The sequence is that of an antimicrobial protein which can
CC CC	be used to control microbial infestations in plants and mammalian
XX XX	animals.
XX XQ	Sequence 590 AA;

Query Match	44.0%	Score 109;	DB 19;	Length 590;
Best Local Similarity	50.0%;	Pred. No. 0.00014;		
Matches 20; Conservative	8;	Mismatches 10;	Indels 2;	Gaps 1;

Qy 3 EDPOTEGCCCCRRCRGQESDPKQOYCRCKEICEEENE 42  
||| :||: |||| || |||| : ||: :  
Db 81 edpqrryeecqecrqee--rqpqcqqrclkrfbegq 118

RESULT	7
W62841	
ID	W62841 standard; Protein; 28 AA
XX	
AC	W62841;

XX	27-OCT-1998	(first entry)
DT		
XX	Stenocarpus sinuatus antimicrobial protein.	
XX		
DE	antimicrobial protein; infestation; control.	
XX		
RW		
XX		
OS	Stenocarpus sinuatus.	
XX		
PN	W09827805-A1.	
XX		
PD	02-JUL-1998.	
XX		
PF	22-DEC-1997;	97WO-AU00874.
XX		
PR	20-DEC-1996;	96AU-0004275.
XX		
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	
XX		
PI	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;	
XX		
DR	WPI; 1998-377279/32.	
XX		
PT	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
XX	useful for controlling microbial infestations of plants or mammals	
PS	Claim 1; Page 66; 96pp; English.	
XX		
CC	The sequence is that of an antimicrobial protein which can	
XX	be used to control microbial infestations in plants and mammalian	
CC	animals.	
XX		
Sequence	28:AA:	
50		

	Query Match	Similarity	63.0%	Score	96:	DB	19:	Length	28:
	Best Local	Conservative	3:	Mismatches	7:	Indels	0:	Gaps	0:
OY	4	DPTCCGCGCQRRCROESDPDROQTCCQ	30						
Db	2	dplrqgqlcmrcqgqekdprdqgqck	28						

RESULT	8	
ID	W62835	
XX	W62835 standard; Protein; 593 AA.	
AC	W62835;	
DT	27-OCT-1998 (first entry)	
DE	Zea mays antimicrobial protein.	
XX	antimicrobial protein; infestation; control.	
XX	Zea mays.	
OS		
FN	W09827805-A1.	
PD	02-JUL-1998.	
PF	22-DEC-1997; 97WO-AU00874.	
PR	20-DEC-1996; 96AU-0004275.	
PA	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	
XX		
XX	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;	
XX	WPI; 1998-377279/32.	
XX	Novel anti-microbial protein from e.g. Macadamia integrifolia -	
DT		

PT useful for controlling microbial infestations of plants or mammals  
PS  
xx Claim 1, Page 58-60, 96pp; English.  
xx  
CC The sequence is that of an antimicrobial protein which can  
CC be used to control microbial infestations in plants and mammalian  
CC animals.  
xx  
SQ Sequence 593 AA;

Query Match	27.68;	Score 68.5;	DB 19;	Length 593;
Best Local Similarity	35.3%;	Pred. No. 3.4;		
Matches .12;	Conservative 10;	Mismatches 11;	Indels 1;	Gaps 1

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Oy      2 QEDPQTECQCQRC-RQESDPQQQCYCQRCK 34
       |: : : |: : |: : |: : |: : |: :
Db    557 eeersgrgecrqlrrhhegqpwetgecmrrcr 590

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RESULT	9
R21079	
ID	R21079 standard; Peptide; 35 AA.

DT 09-APR-1992 (first entry)

DE Antimicrobial maize peptide, CMIII.

KW Maize; CMIIT; corn; pathogen.

05 Zea mays.

PN EP465009-A.

PD 08-JAN-1992.

PF 05-JUN-1991; 91EP-0305064

PR 05-JUN-1990; 90US-0536127.

PA (PION-) PIONEER HI-BRED INT.

PI Duvick JP, Rood TA, Rao AG;

DR WPI; 1992-010214/02.

PT Use of maize seed peptide CMIII and DNA encoding it - for killing  
PT or inhibiting plant pathogenic microorganisms.

PS Example 2; Page 5; 21pp; English.

CC The peptide (SEQ ID NO 1) was purified from public corn variety B73  
CC and Proprietary corn variety M161. It is basic and has a total  
CC mol. wt. of 3900 daltons. The peptide sequence was used to design  
CC probes which were used to screen a maize genomic or cDNA library.  
CC The isolated CMII gene can be used to prepare an expression vector  
CC for prodn. of recombinant CMII for use in controlling plant patho-  
CC genic organisms.  
CC See also Q20272 and 3.

SQ Sequence 35 AA;

Query Match	26.8%;	Score 66.5;	DB 13;	Length 35;
Best Local Similarity	44.0%;	Pred. No. 0.4;		
Matches 11;	Conservative 6;	Mismatches 7;	Indels 1;	Gaps 1;

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OY      11 QCQRC_RQGSDPRQQCYCQRCK 34  
          :|:|:| |: | : | | ||:  
Db      6 ecrqqlrhhegpyetqcmrrc 30
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RESULT 10
W62837
ID W62837 standard; Protein; 637 AA.
XX
AC W62837;
XX
DT 27-OCT-1998 (first entry)
XX
DE Hordeum vulgare antimicrobial protein.
XX
KW antimicrobial protein; infestation; control.
XX
OS Hordeum vulgare.
XX
PN W62837;
XX
PD 02-JUL-1998.
XX
PE 22-DEC-1997; 97MO-AU00874.
XX
PR 20-DEC-1996; 96AU-0004275.
XX
PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
XX
PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX
DR WPI; 1998-377279/32.
XX
PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX
PS useful for controlling microbial infestations of plants or mammals
XX
PS Claim 1; Page 60-62; 96pp; English.
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 637 AA;

Query Match 26.8%; Score 66.5; DB 19; Length 637;
Best Local Similarity 39.4%; Pred. No. 6.1;
Matches 13; Conservative 7; Mismatches 8; Indels 5; Gaps 2;

QY 10 QOCRCRCQOESDPROQOYQCRCKEICEEE 42
   |||:||||: || | ||:|:|:|
Db 42 qgcvggrgrgr--pr---ysharcvgecrddq 69

RESULT 11
W62836
ID W62836 standard; Protein; 33 AA.
XX
AC W62836;
XX
DT 27-OCT-1998 (first entry)
XX
DE Zea mays antimicrobial protein.
XX
KW antimicrobial protein; infestation; control.
XX
OS Zea mays.
XX
PN W62836;
XX
PD 02-JUL-1998.
XX
PE 22-DEC-1997; 97MO-AU00874.
XX
PR 20-DEC-1996; 96AU-0004275.
XX
PA (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.

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XX
PI Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
XX
DR WPI; 1998-377279/32.
XX
PT Novel anti-microbial protein from e.g. Macadamia integrifolia -
XX
PS useful for controlling microbial infestations of plants or mammals
XX
PS Disclosure; Page 60; 96pp; English.
XX
CC The sequence is that of an antimicrobial protein which can
CC be used to control microbial infestations in plants and mammalian
CC animals.
XX
SQ Sequence 33 AA;

Query Match 26.4%; Score 65.5; DB 19; Length 33;
Best Local Similarity 44.0%; Pred. No. 0.49;
Matches 11; Conservative 6; Mismatches 7; Indels 1; Gaps 1;

QY 11 QOCRCRCQOESDPROQOYQCRCK 34
   ||:| | | | | | | | | | | | |
Db 6 ecrrgcrlrhngqpwetqecmrrr 30

RESULT 12
Y22191
ID Y22191 standard; Protein; 910 AA.
XX
AC Y22191;
XX
DT 10-SEP-1999 (first entry)
XX
DE Mouse brain CNG-1 protein sequence.
XX
KW BCNG; brain cyclic nucleotide gated ion channel; epilepsy; hyperalgesia;
KW Alzheimer's Disease; Parkinson's Disease; long QT syndrome; dyslexia;
KW sick sinus syndrome; age-related memory loss; cystic fibrosis;
KW sudden death syndrome; pacemaker rhythm dysfunction; sensory disorder;
KW auditory disorder; respiratory disorder; attention deficit disorder;
KW learning disability; drug addiction; therapy; mBCNG-1.
XX
OS Mus sp.
XX
PN W62836;
XX
PD 01-JUL-1999.
XX
PE 23-DEC-1998; 98MO-US27630.
XX
PR 28-MAY-1998; 98US-0086436.
XX
PR 23-DEC-1997; 97US-0997685.
XX
PA (UYCO ) UNIV COLUMBIA NEW YORK.
XX
PI Bartsch D, Grant S, Kandel ER, Santoro B, Siegelbaum S;
XX
PI Tibbs G;
XX
DR WPI; 1999-418922/35.
XX
DR N-PSDB; X84442.
XX
PT An isolated nucleic acid encoding a brain or heart cyclic
XX
PT nucleotide-gated ion channel
XX
PS Claim 16; Page 185-188; 213pp; English.
XX
CC This sequence is the brain cyclic nucleotide-gated ion channel
XX
CC (BCNG) of the invention, designated mBCNG-1. BCNG and h-KNG-related
XX
CC proteins are useful in screening for compounds that modulate, interact or
XX
CC affect expression. Compounds, e.g. antagonists and agonists, identified
XX
CC in the methods are useful for modulating BCNG or BCNG-related protein
XX
CC activity. Modulation is increased or decreased ion permissivity or ion

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